

1 **A**

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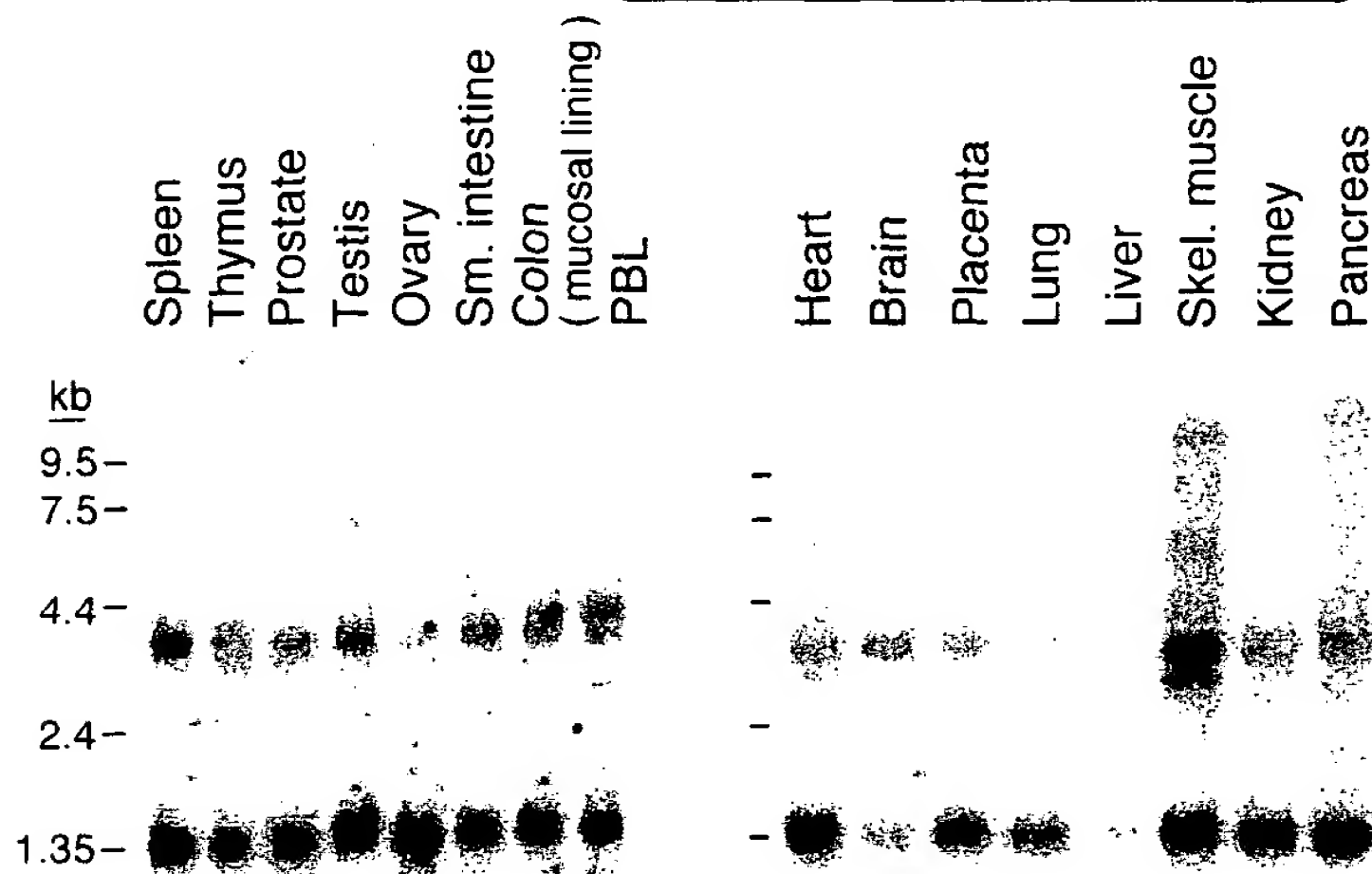
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M P K R K V T F Q G V G D E E D E D E I
61  ATTGTCCCCAAGAAGAAGCTGGTGGACCCTGTGGCTGGGTCAGGGGGTCCTGGGAGCCGC
I V P K K K L V D P V A G S G G P G S R
121 TTTAAAGGCAAACACTCTTTGGATAGCGATGAGGAGGAGGATGATGATGATGGGGGGTCC
F K G K H S L D S D E E E D D D G G S
181 AGCAAATATGACATCTTGGCCTCAGAGGATGTAGAAGGTCAGGAGGCAGCCACACTCCCC
S K Y D I L A S E D V E G Q E A A T L P
241 AGCGAGGGGGGTGTTCCGATCACACCCTTTAACCTGCAGGAGGAGATGGAGGAAGGCCAC
S E G G V R I T P F N L Q E E M E E G H
301 TTTGATGCCGATGGCAACTACTTCTGAACCGGGATGCTCAGATCCGAGACAGCTGGCTG
F D A D G N Y F L N R D A Q I R D S W L
361 GACAACATTGACTGGGTGAAGATCCGGGAGCGGCCACCTGGCCAGCGCCAGGCCTCAGAC
D N I D W V K I R E R P P G Q R Q A S D
421 TCGGAGGAGGAGGACAGCTTGGGCCAGACCTCAATGAGTGCCCAAGCCCTCTGGAGGGA
S E E E D S L G Q T S M S A Q A L L E G
481 CTTTGGAGCTCCTATTGCCTAGAGAGACAGTGGCTGGGGCACTGAGGCGTCTGGGGGCC
L L E L L L P R E T V A G A L R R L G A
541 CGAGGAGGAGGCAAAGGGGAGAAAGGGGCCTGGGCAACCCAGTTCCCTCAGCGCCTGGAC
R G G G K G R K G P G Q P S S P Q R L D
601 CGGCTCTCCGGGTGGCCGACCAGATGGTGGCCCGGGGCAACCTTGGTGTGTACCAGGAA
R L S G L A D Q M V A R G N L G V Y Q E
661 ACAAGGGAACGGTGGCTATGCGTCTGAAGGGTTTGGGGTGTGAGACCCTAGGACCCAC
T R E R L A M R L K G L G C Q T L G P H
721 AATCCACACCCCCACCTCCCTGGACATGTTGCTGAGGAGTTGGCGGAGGAGGAAGT
N P T P P P S L D M F A E E L A E E E L
781 GAGACCCCAACCCCTACCCAGAGAGGAGAAGCAGAGTCGCGGGGAGATGGTCTGGTGGAT
E T P T P T Q R G E A E S R G D G L V D
841 GTGATGTGGGAATATAAGTGGGAGAACACGGGGGATGCCGAGCTGTATGGGCCCTTACC
V M W E Y K W E N T G D A E L Y G P F T
901 AGCGCCAGATGCAGACCTGGGTGAGTGAAGGCTACTTCCCGGACGGTGTATTGCCGG
S A Q M Q T W V S E G Y F P D G V Y C R
961 AAGCTGGACCCCCCTGGTGGTCACTTCTACAACCTCCAAACGCATTGACTTTGACCTCTAC
K L D P P G G Q F Y N S K R I D F D L Y*
1021 ACCTGAGCCTGCTGGGGGCCAGTTTGGTGGGCCCTTCTTCTGGAATTTGTGGAGGAG
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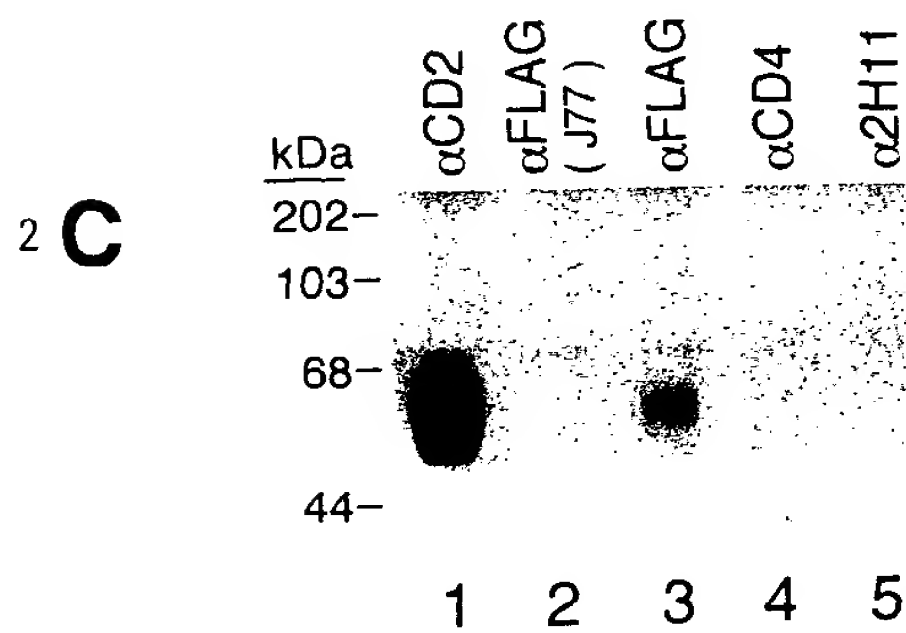
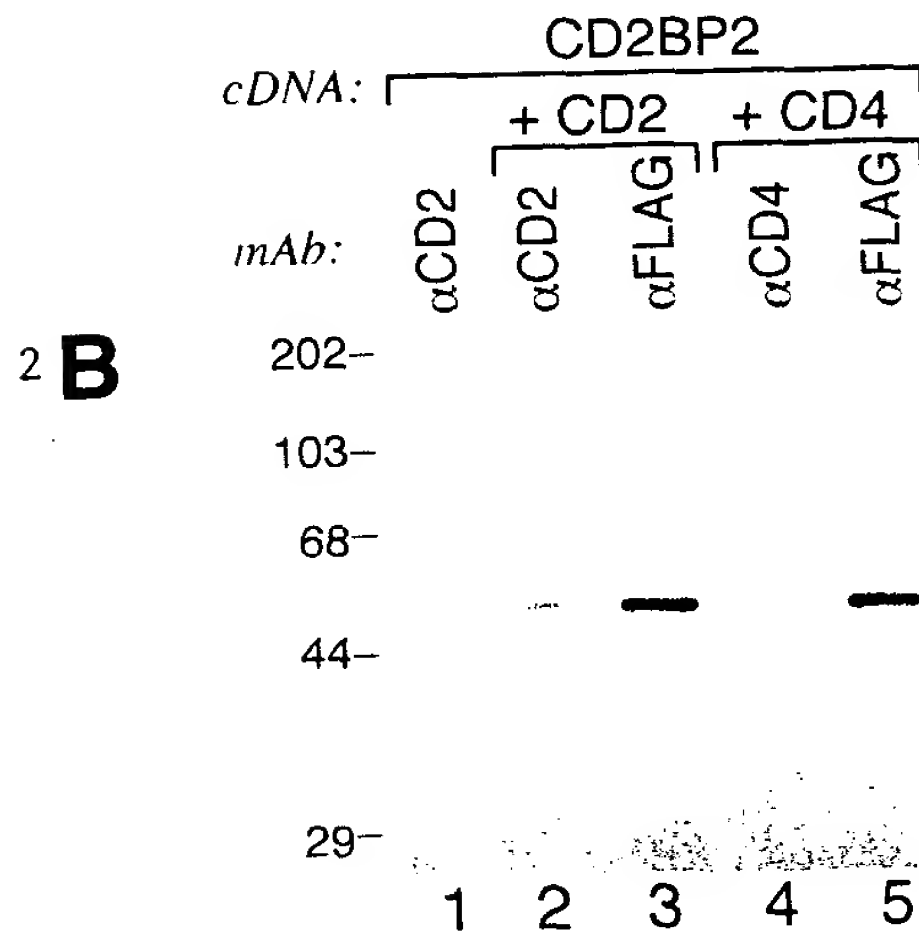
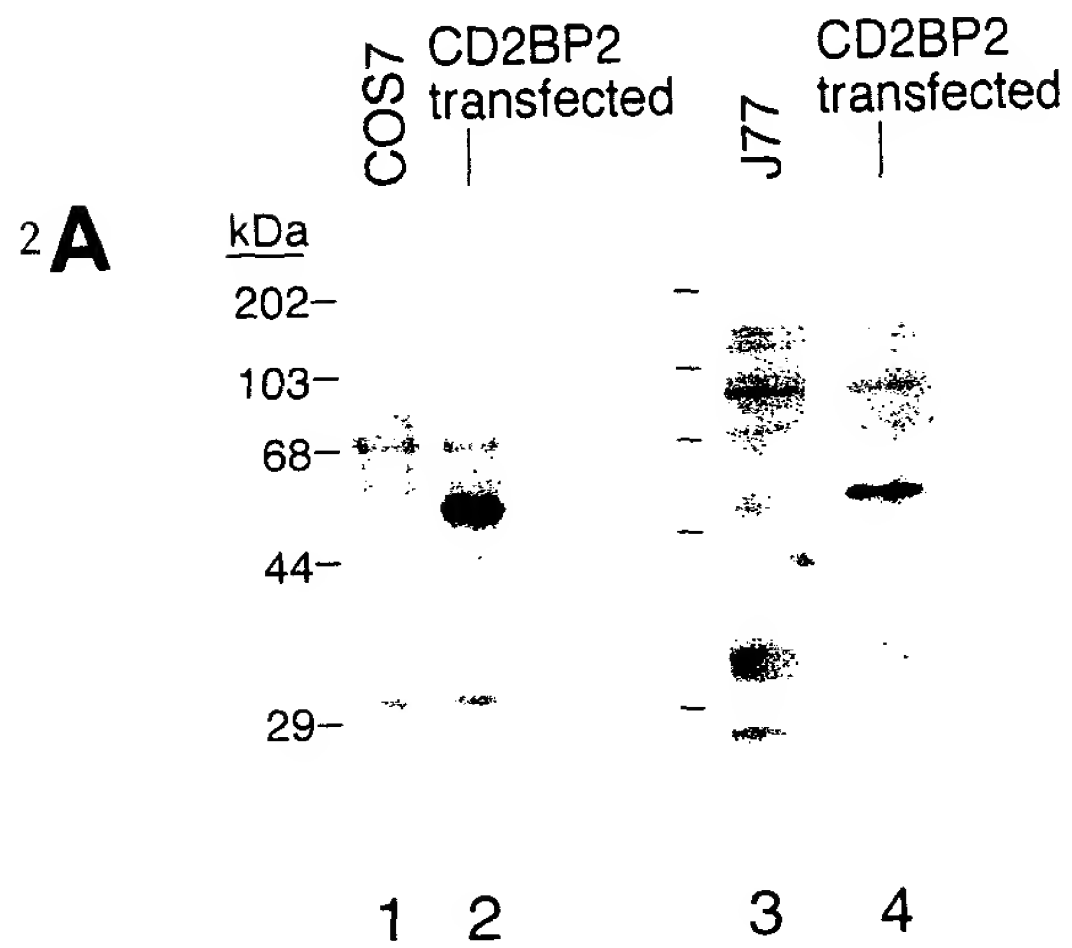
1 **B**

CD2BP2 291-317 G - D A E L Y G P F T S A Q M Q T W V S E G Y F P D G  
*C. el.* (z81109) G P D S E K Y G P Y M S K D M L F W L Q A G Y F N D G  
*C. el.* (u23147) D P T E T R R G P F P K D Q M N V W F K A G Y F T D E  
*C. el.* (p34520) D D R G T V Q G P Y G A S T V L D W Y Q K G Y F S D N  
*S. ce.* (SMY2) D T Q G Q I H G P F T T Q M M S Q W Y I G G Y F A S T  
*S. ce.* (s61962) D S N G N I Q G P F G T N N M S Q W Y Q G G Y F T P T

G P F	M	W	G Y F
Y	V		

1 **C**





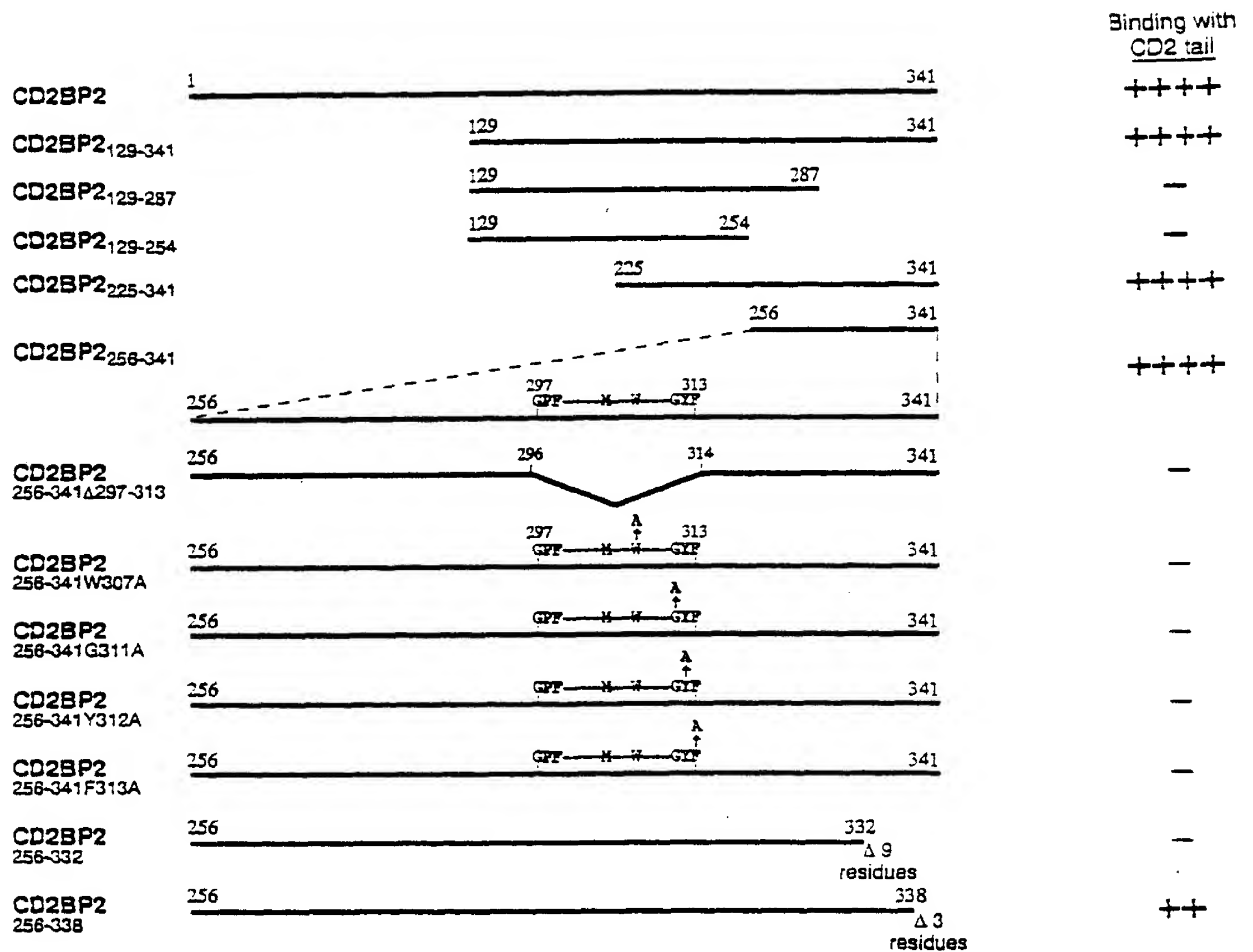


FIG. 3A

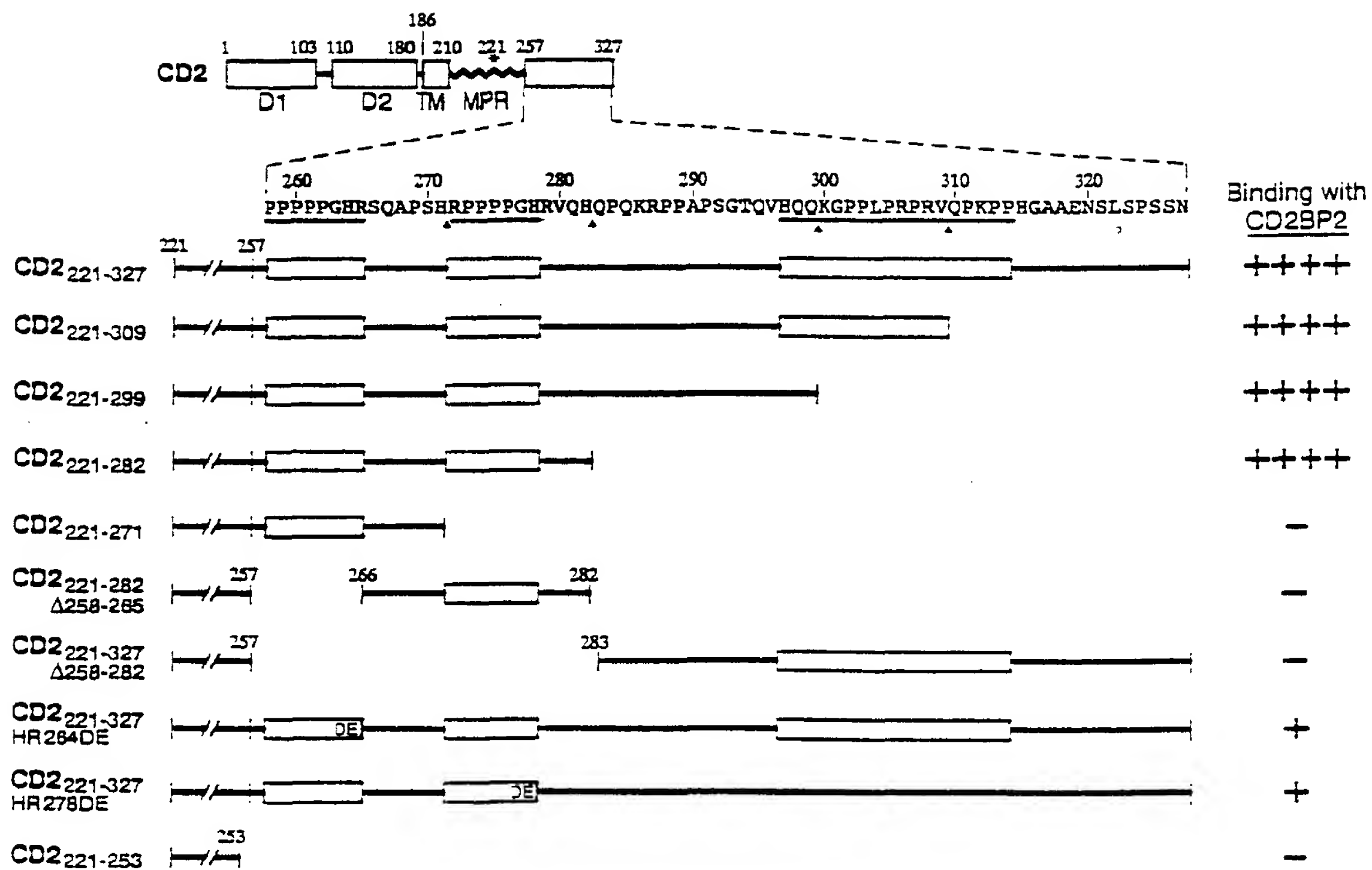
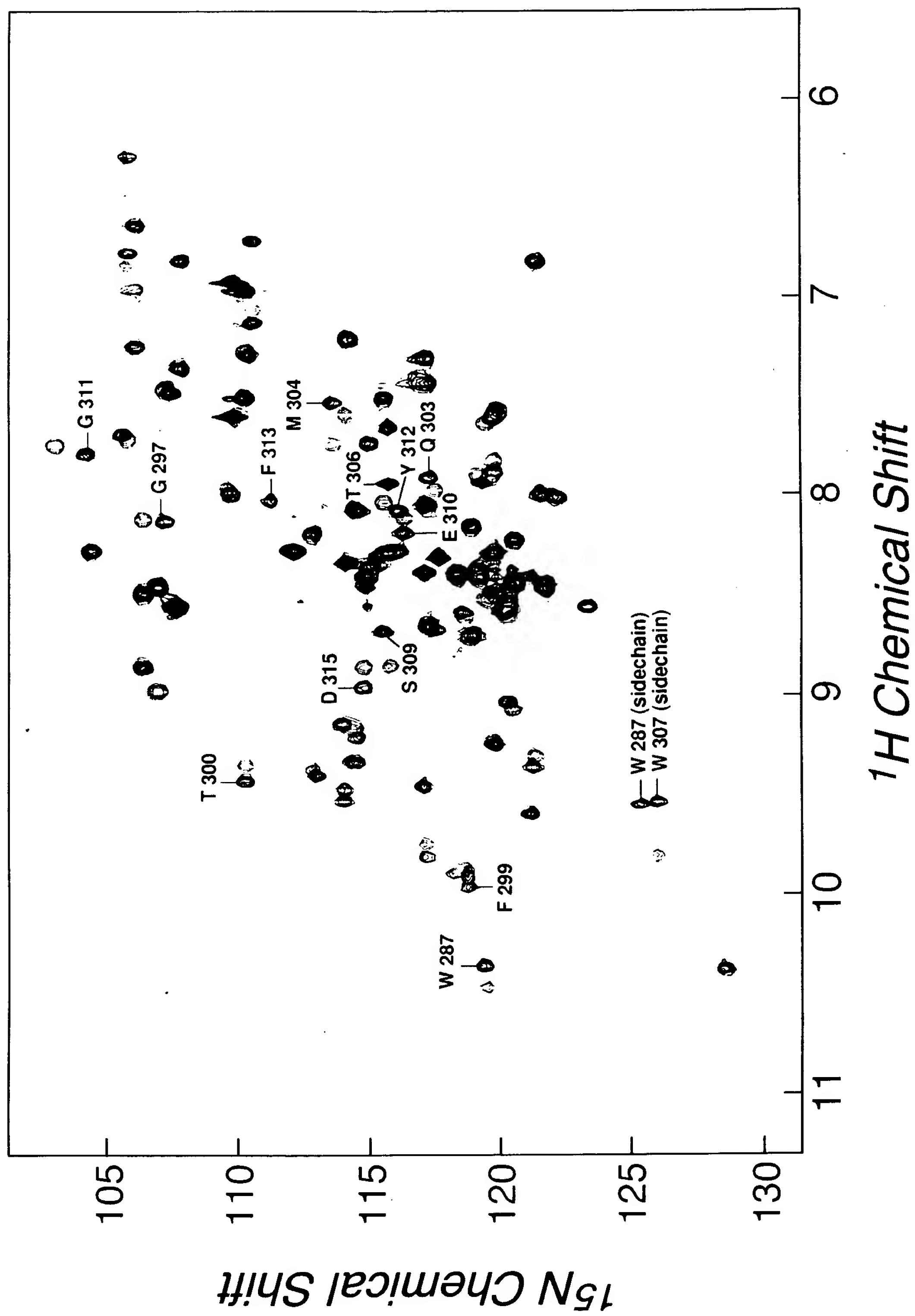


FIG. 3B



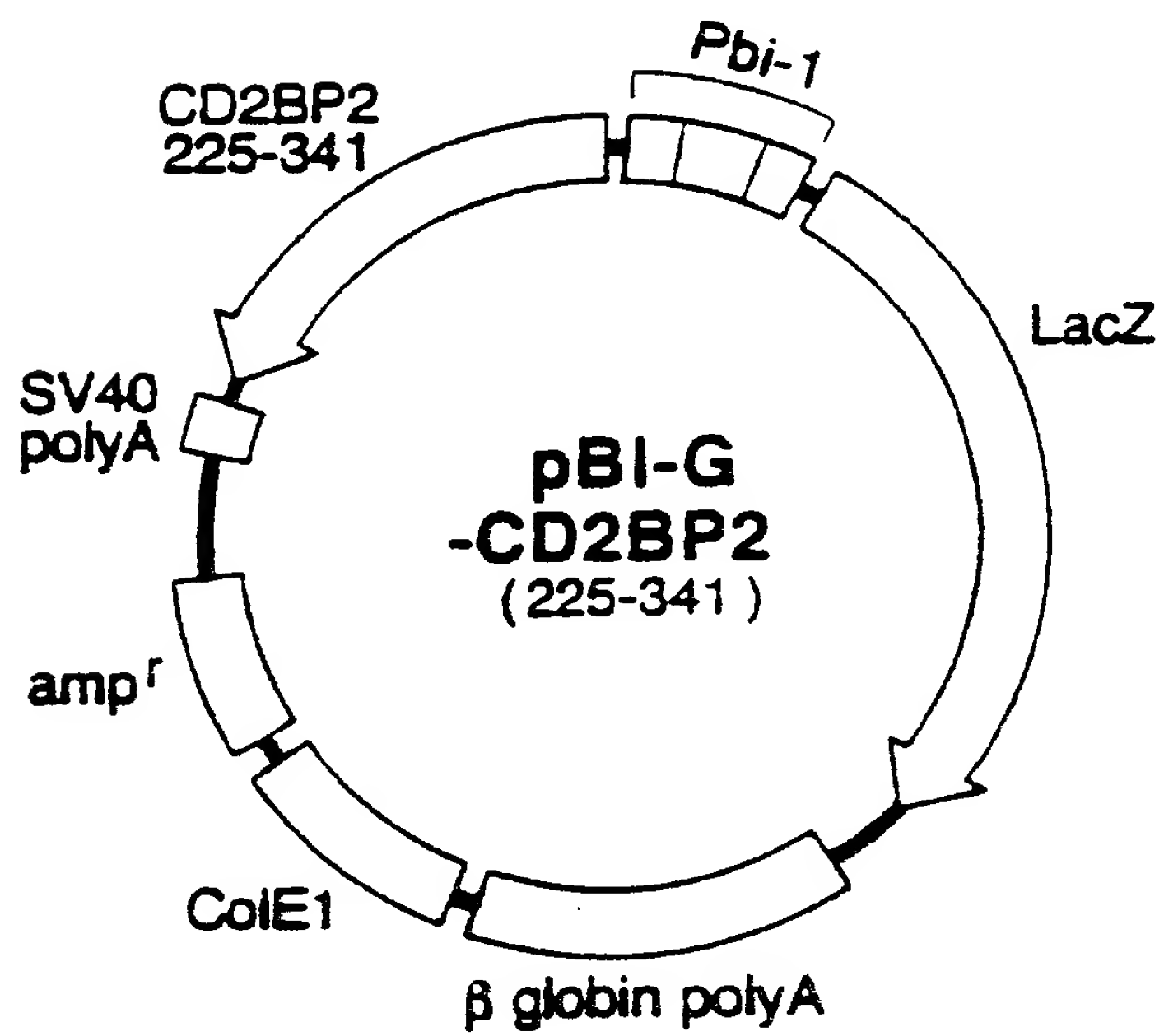


FIG. 5A

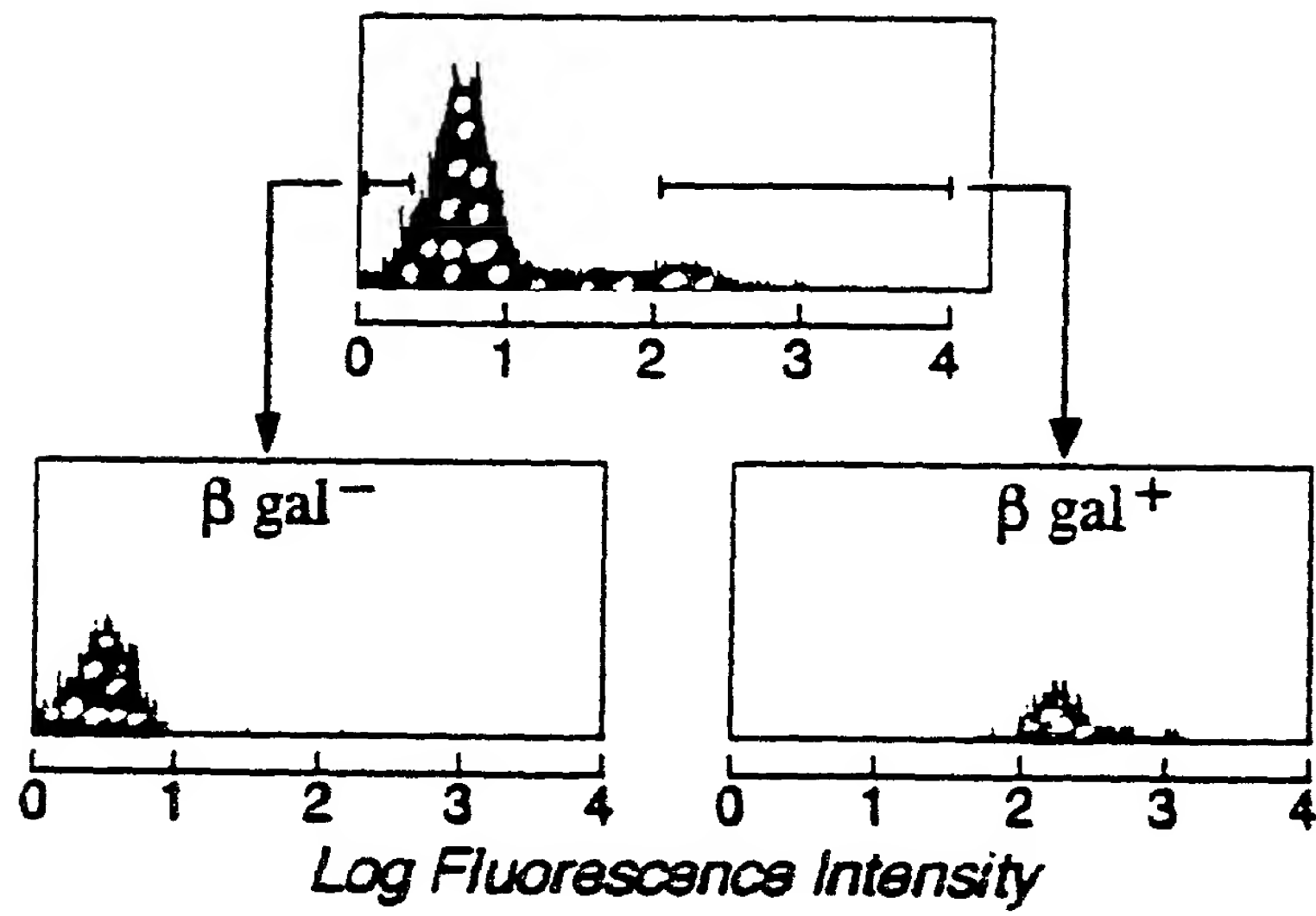


FIG. 5B

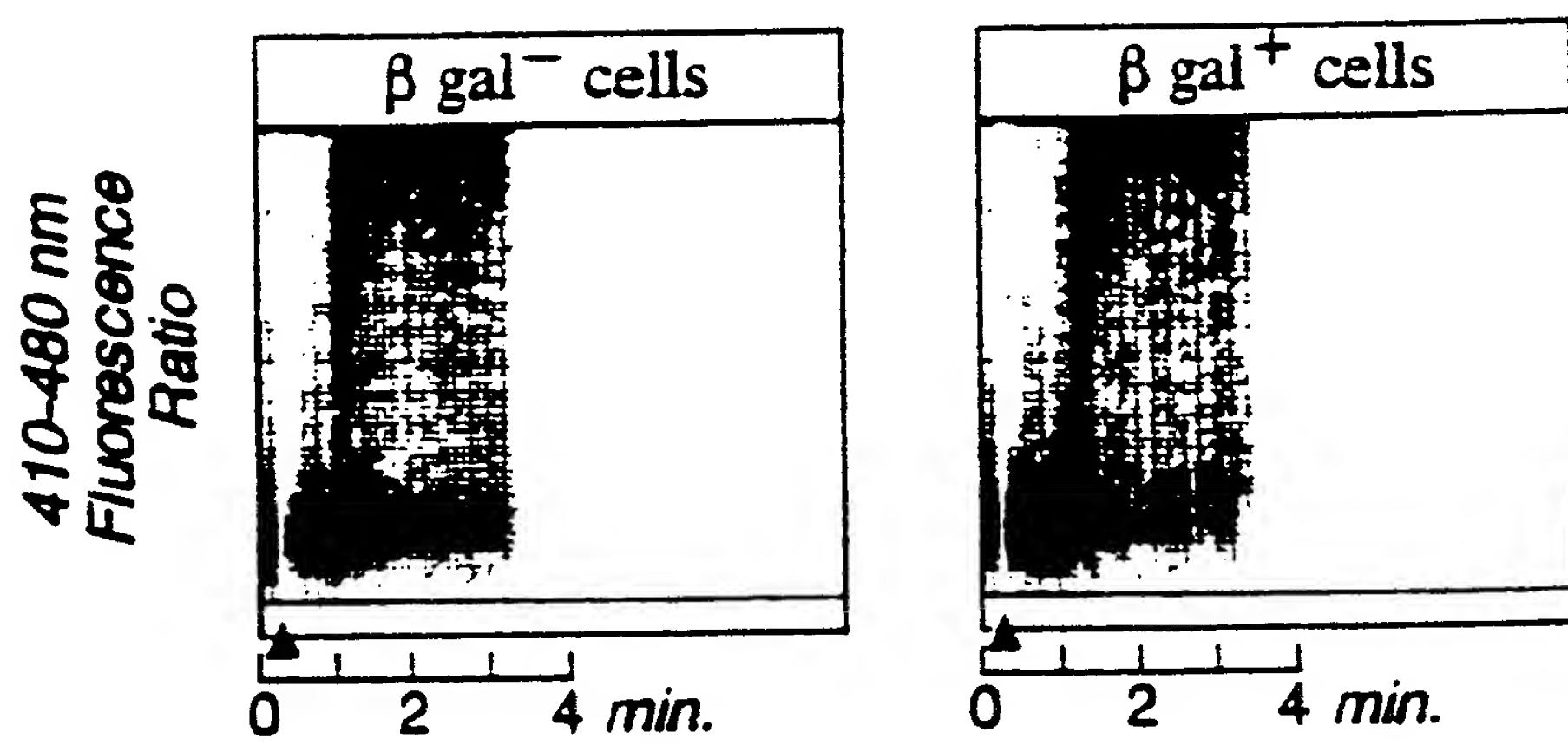


FIG. 5C



FIG. 5D

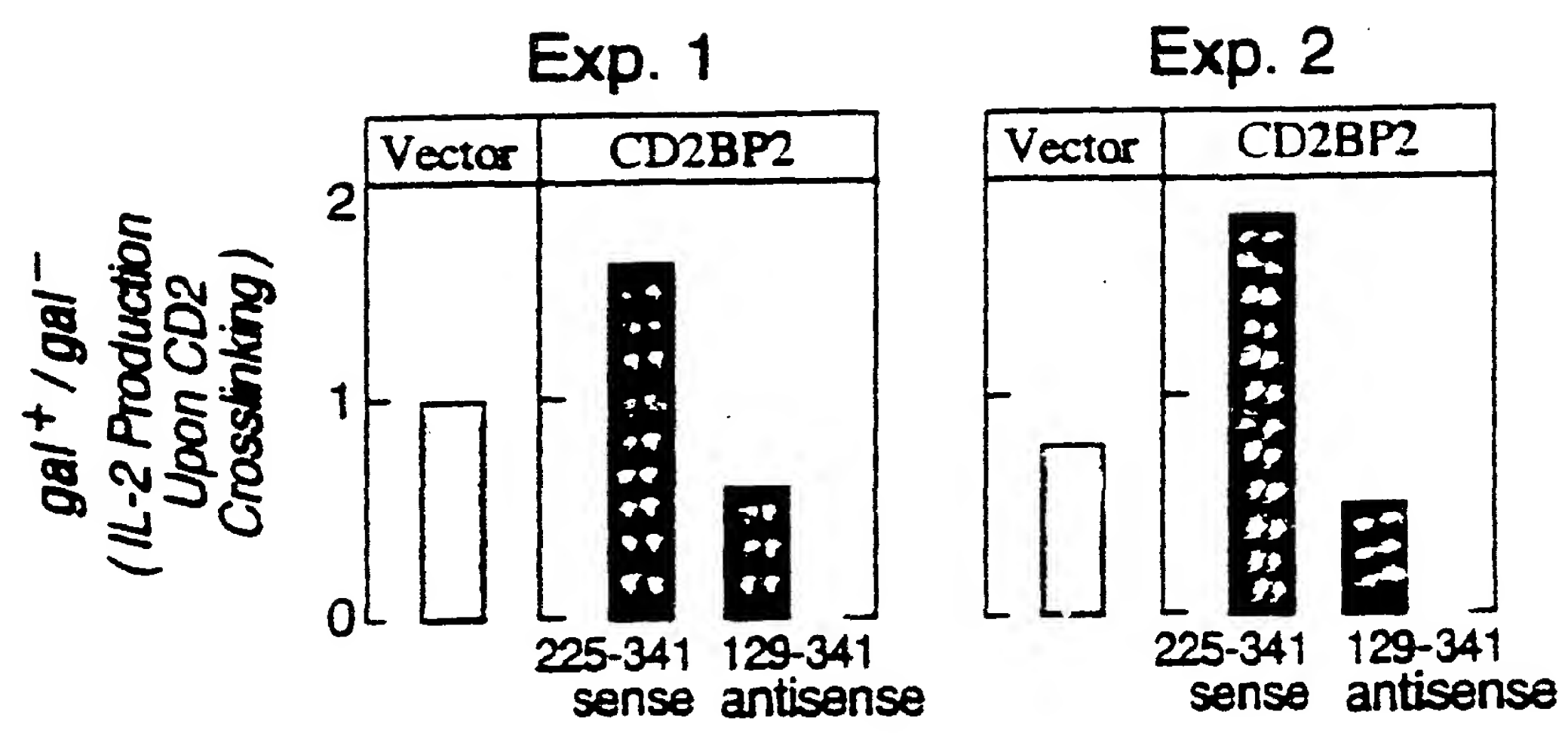


FIG. 5D

107030-907E600

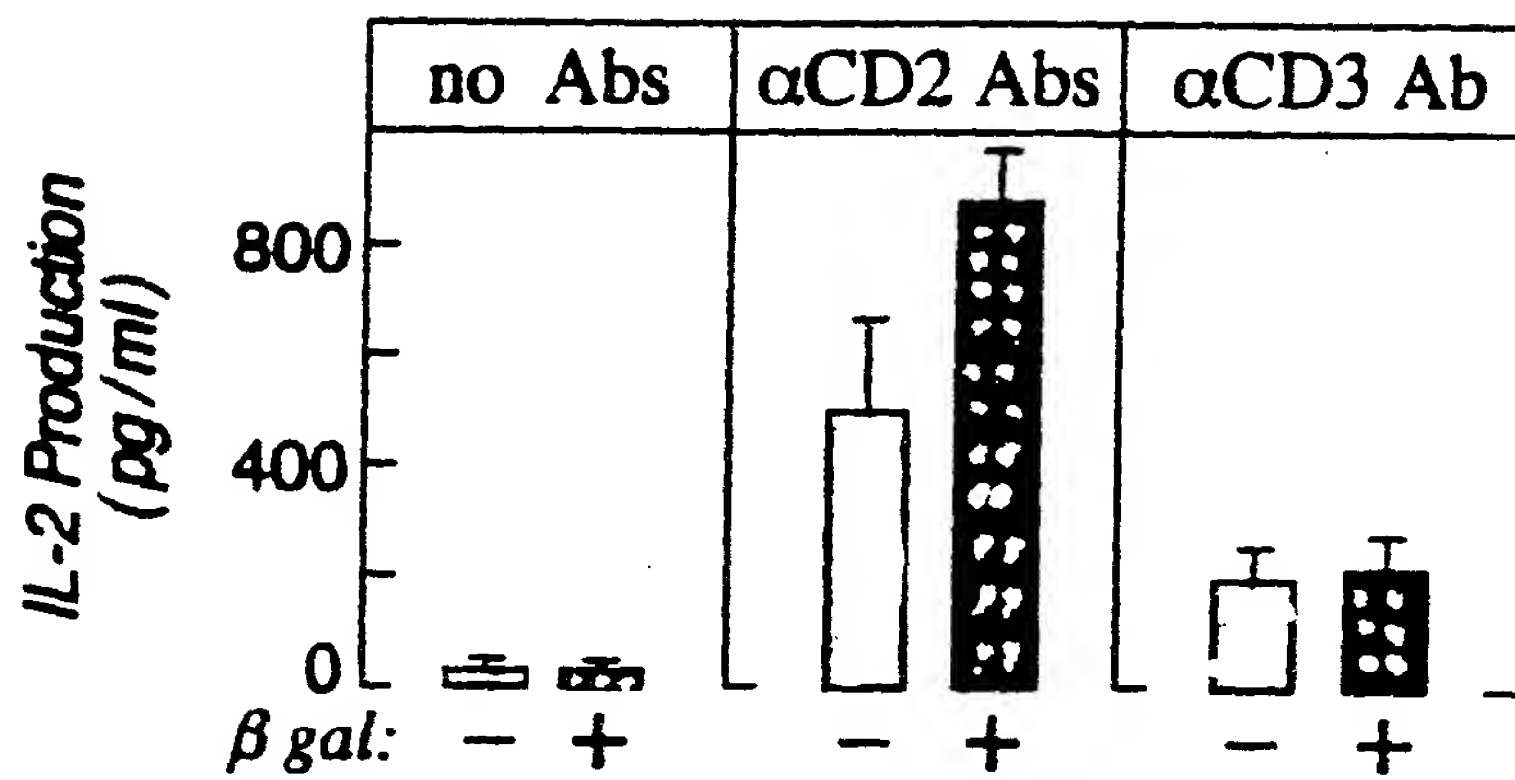


FIG. 5E

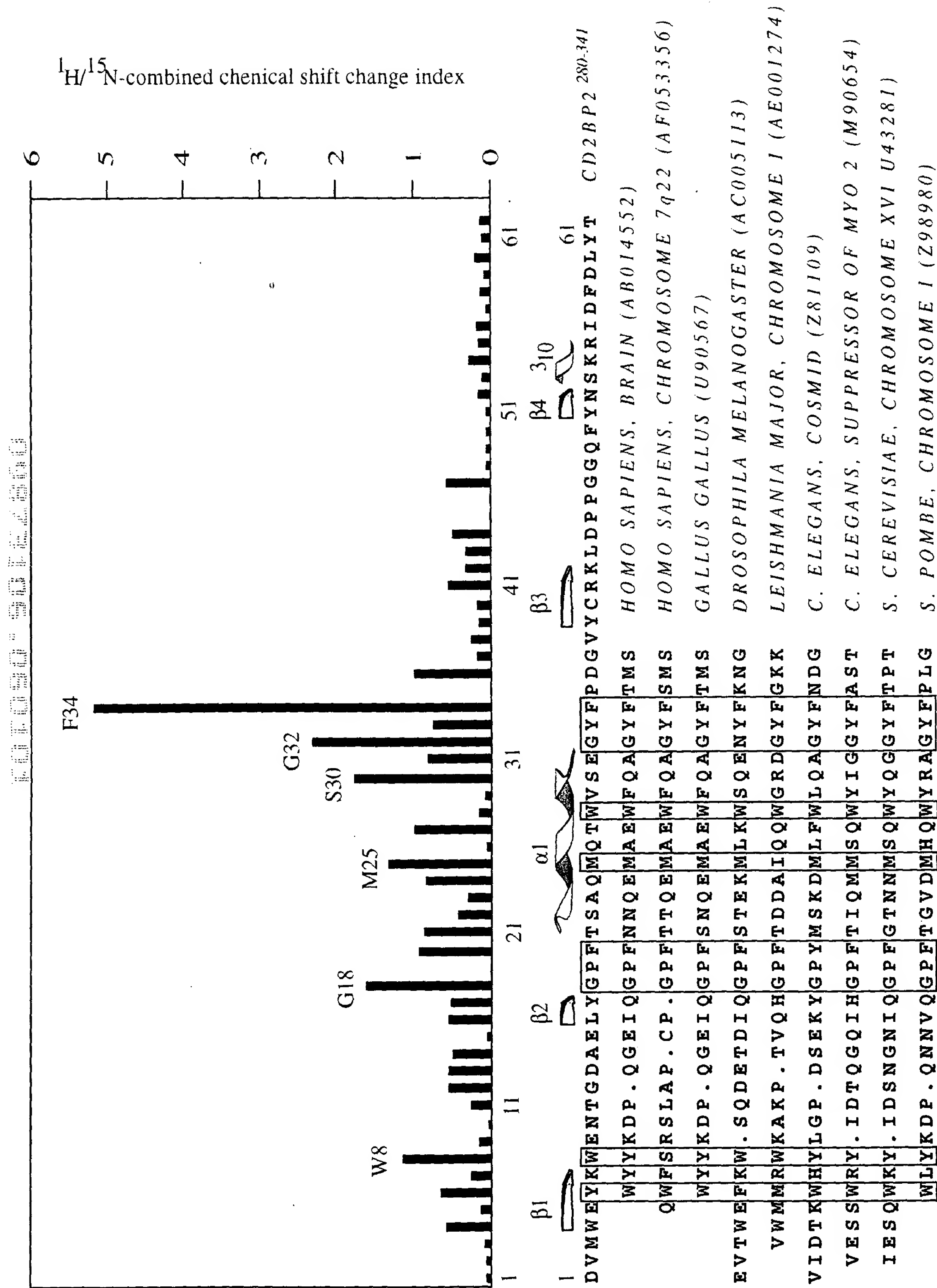


Fig. 6

FIG. 7A

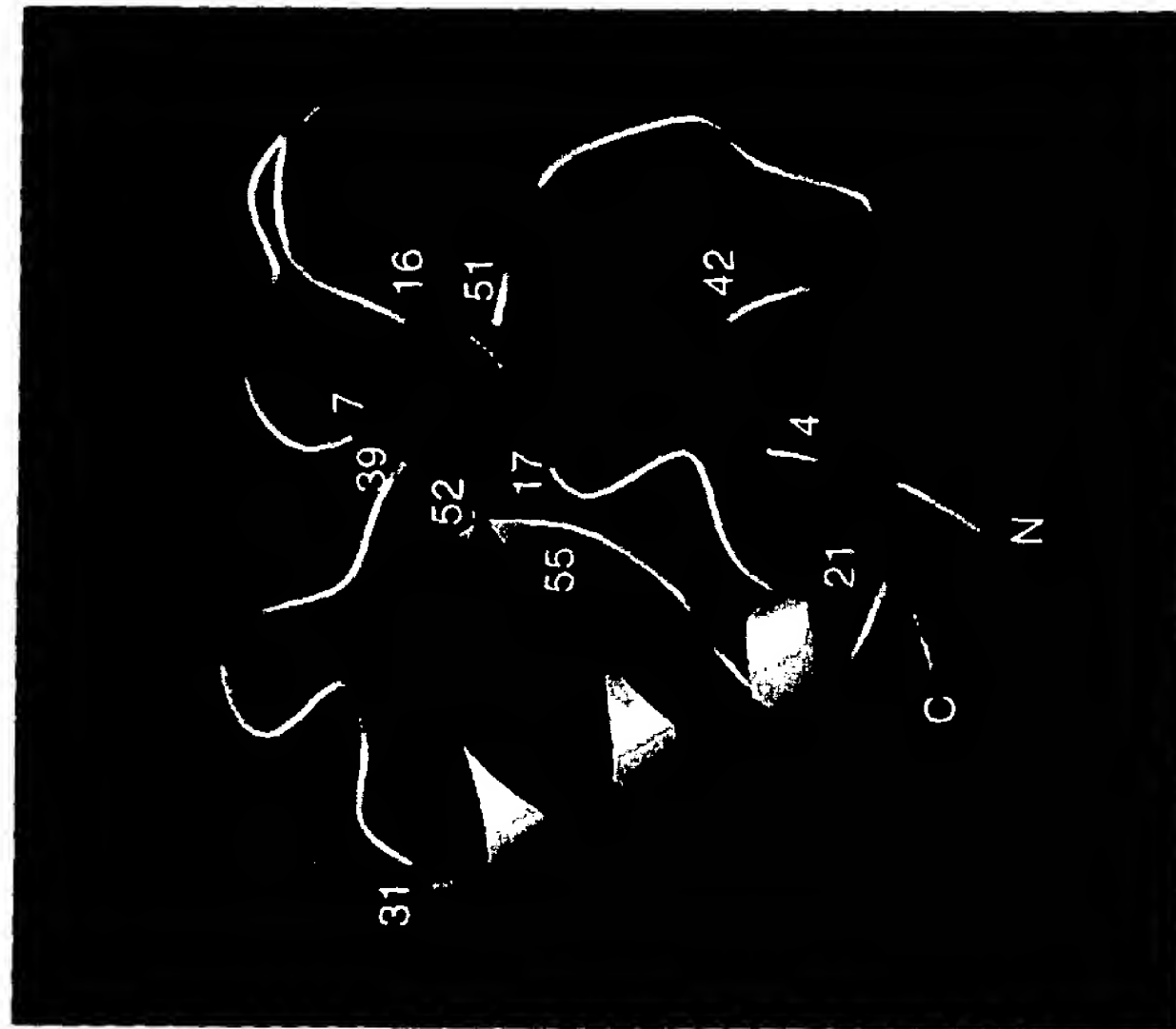


Fig. 7A

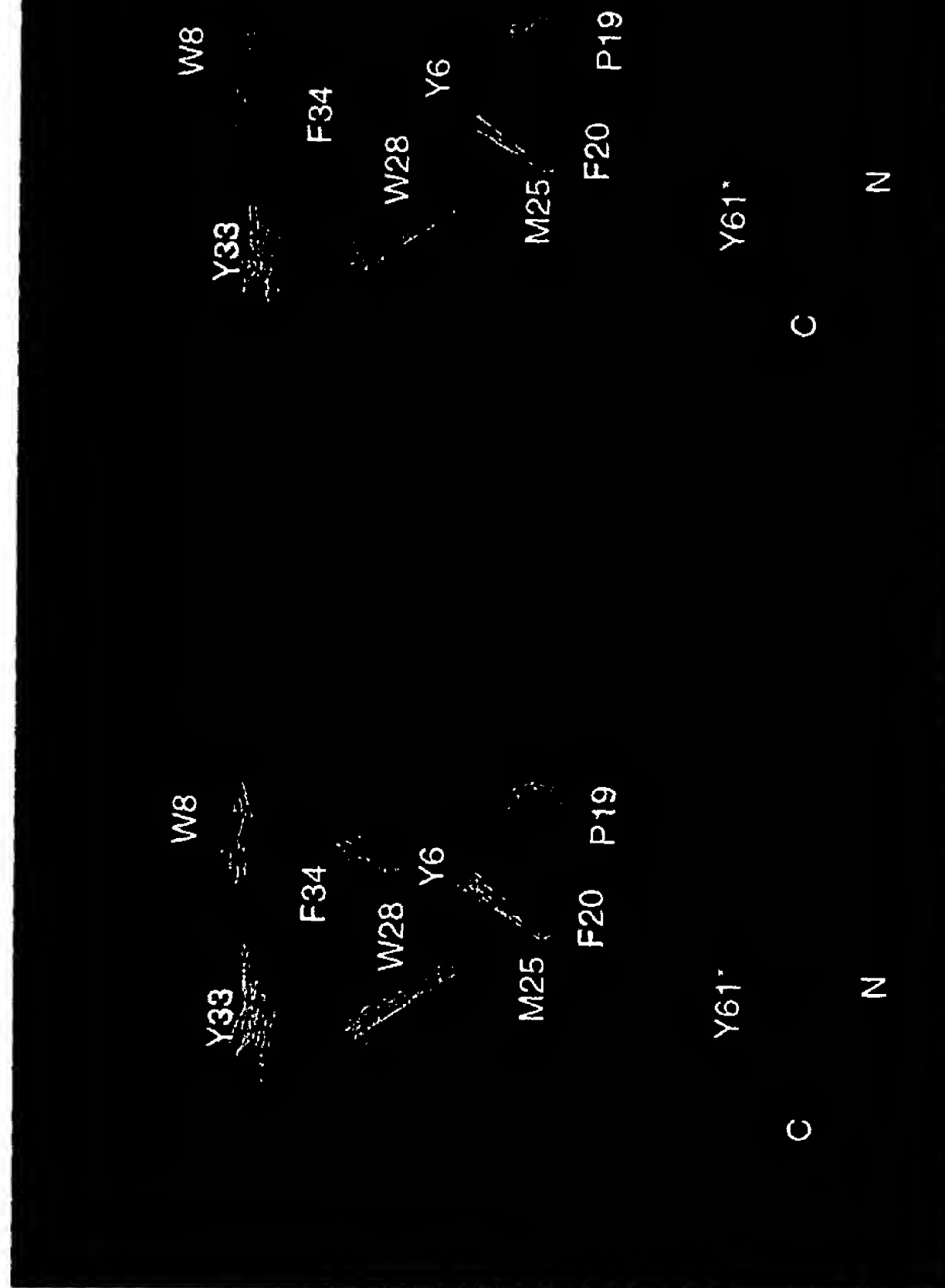


Fig. 7B

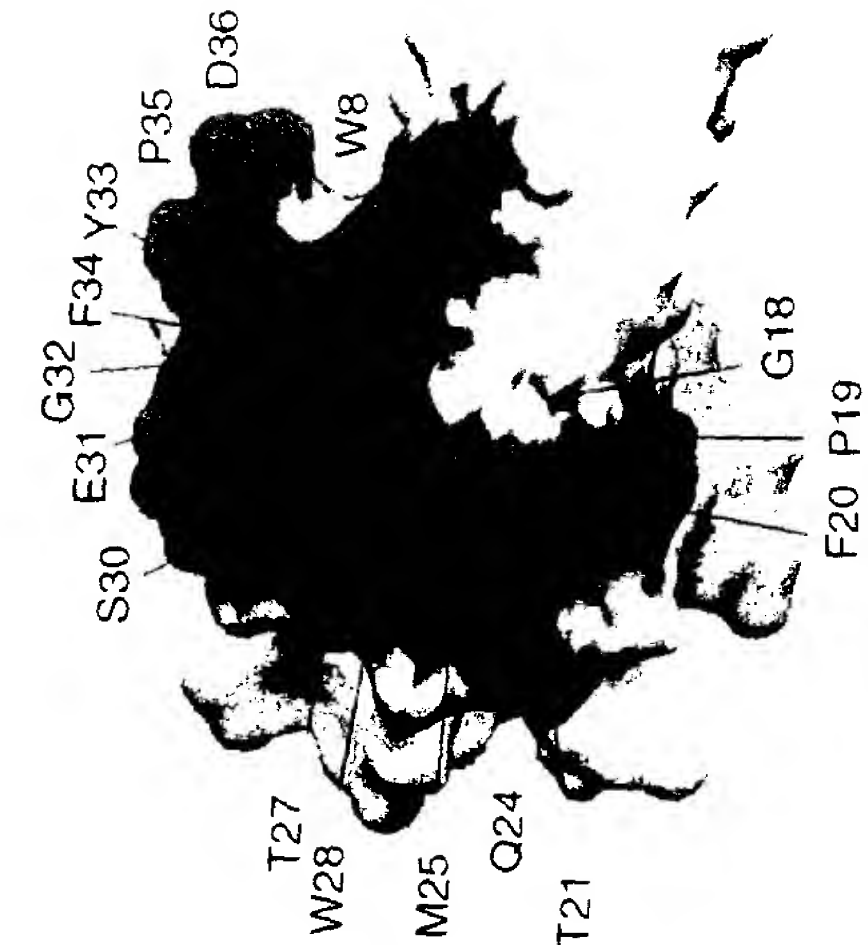


Fig. 8A

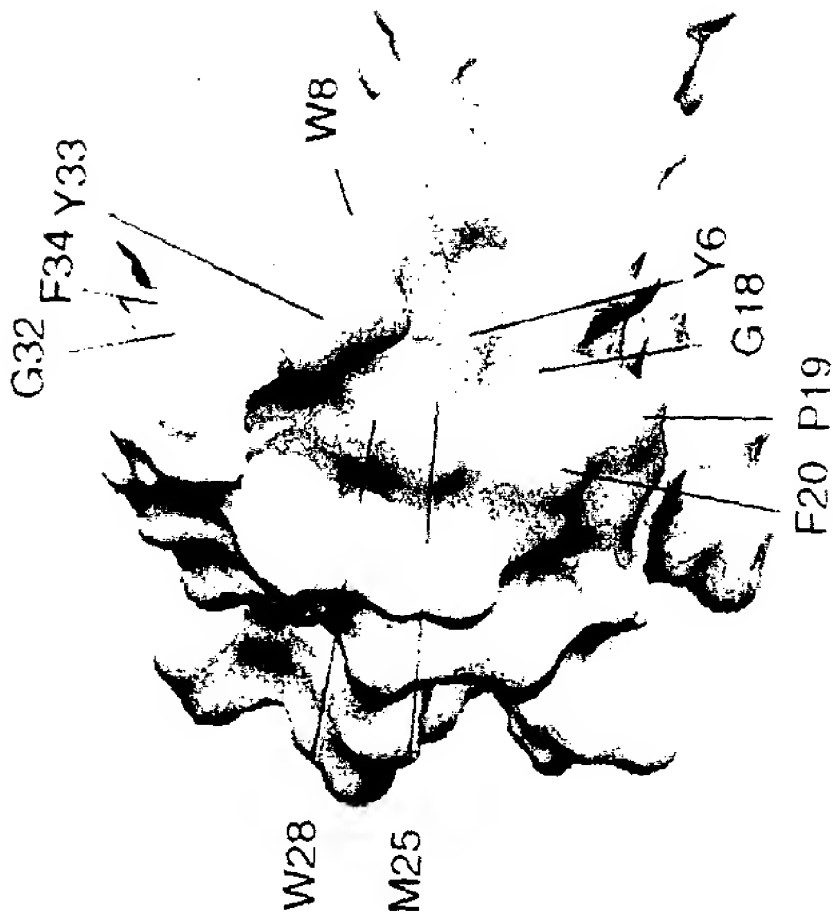


Fig. 8B

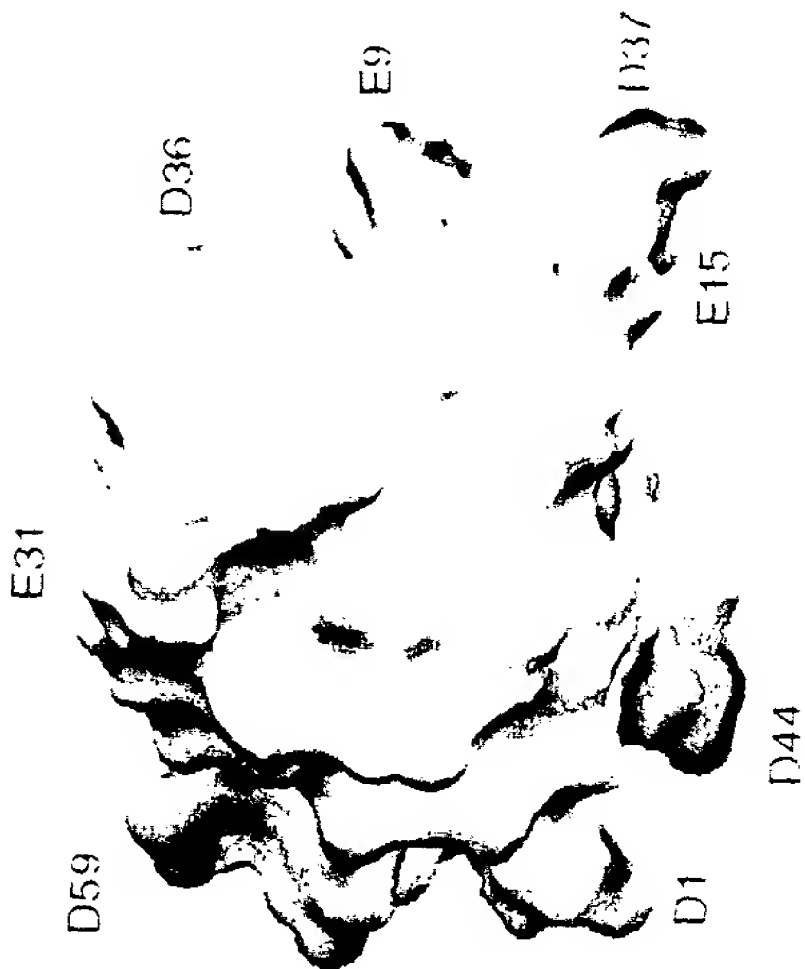


Fig. 8C